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**INFORMATION DISCLOSURE
STATEMENT BY APPLICANT**

(use as many sheets as necessary)

Sheet 1

of

2

Complete If Known

Application Number	09/848,866
Filing Date	May 4, 2001
First Named Inventor	Duncan McRee et al.
Group Art Unit	2171
Examiner Name	Borin, Michael L.
Attorney Docket Number	SYRTECH-5001-U

U.S. PATENT DOCUMENTS

Examiner Initials ¹	Cite No. ¹	Document Number	Publication Date/ Issue Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
		Number - Kind Code ² (if known)			
MM	AA	US- 5,375,207	12-20-1994	Blakely et al.	
	AB	US- 5,436,850	07-25-1995	Eisenberg et al.	
	AC	US- 5,600,833	02-04-1997	Senn et al.	
	AD	US- 5,859,972	01-12-1999	Subramaniam et al.	
	AE	US- 5,878,373	03-02-1999	Cohen et al.	
MM	AF	US- 6,240,374B1	05-29-2001	Cramer et al.	

FOREIGN PATENT DOCUMENTS

Examiner Initials ¹	Cite No. ¹	Foreign Patent Document	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear	T ⁶
		Country Code ³ - Number ⁴ - Kind Code ⁵ (if known)				
MB	AG	WO 01/09595 A2-A3	02-08-2001	Emerald Biostructures, Inc.		
MB	AH	WO 03/032558 A2-A3	04-17-2003	Emerald Biostructures, Inc.		

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Considered

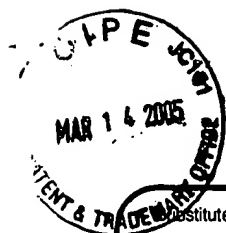
04/2005

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Sheet 2 of 2	Attorney Docket Number	SYRTECH-5001-U	

OTHER PRIOR ART – NON PATENT LITERATURE DOCUMENTS			
Examiner Initials *	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T ²
MB	AI	BERMAN, Helen et al., The Protein Data Bank, Nucleic Acids Research, 2000, Vol. 28, No. 1, 235-242	
	AJ	FOADI, J. et al., A flexible and efficient procedure for the solution and phase refinement of protein structures, Acta Cryst. (2000), D56, 1137-1147	
	AK	GRAY, Peter M.D. et al., Macromolecular structure information and databases, TIBS 21 (July 1996), pp. 251-254	
	AL	HOLM, Liisa et al., Protein Structure Comparison by Alignment of Distance Matrices, J. Mol. Biol. (1993) 233, 123-138	
	AM	MOCHALKIN, IGOR et al., "High-Throughput Structure Determination in an Informatics Environment", http://www.accelrys.com/webzine (8/1/02), 4 pages.	
	AN	MURZIN, Alexey G. et al., SCOP: A Structural Classification of Proteins Database for the Investigation of Sequences and Structures, J. Mol. Biol. (1995) 247, 536-540	
	AO	NAVAZA, J., AmoRe: an Automated Package for Molecular Replacement, Acta. Cryst. (1994), A50, 157-163	
MB	AP	ROBINSON, Daniel D., Partial Molecular Alignment via Local Structure Analysis, J. Chem. Inf. Comput. Sci. 2000, 40, 503-512	

Examiner Signature		Date Considered	04/2005
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